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WO 2005/080426

PCT/GB2005/000572

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Figure 1

MAAAASPAFLRLPLLLLSSWCRTGLADPHSLCYDITVIPKIRPGPRWCAVQGQVDE
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Figure 2

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1 cctgcgagcc gccaggtgat ccacgggctg ggcttcgctt ctgctgtccc ctgcgatcca
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121 tgctgtccag ctgggtgcagg accgggctgg ccgaccctca ctctcttgc tatgacatca
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1141 aaa

Figure 3

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121 cctaagttca gacctggacc acgggttgtt gcggttcaag gccaggtgga tgaaaagact
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541 tccttcatt acatctcaat gggagactgc acaggatggc ttgaggactt cttgatggc
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Figure 4

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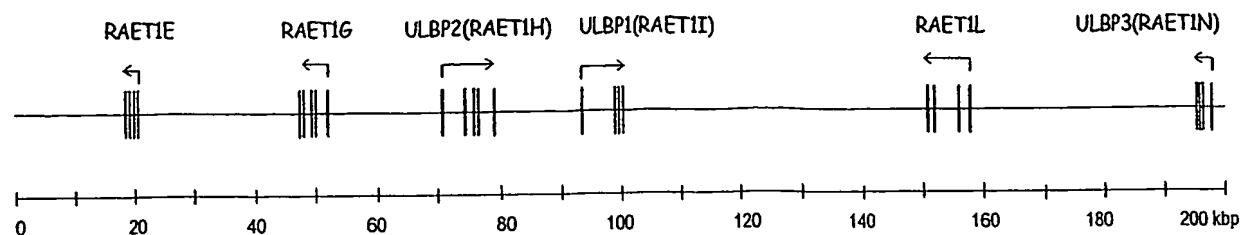


Figure 5

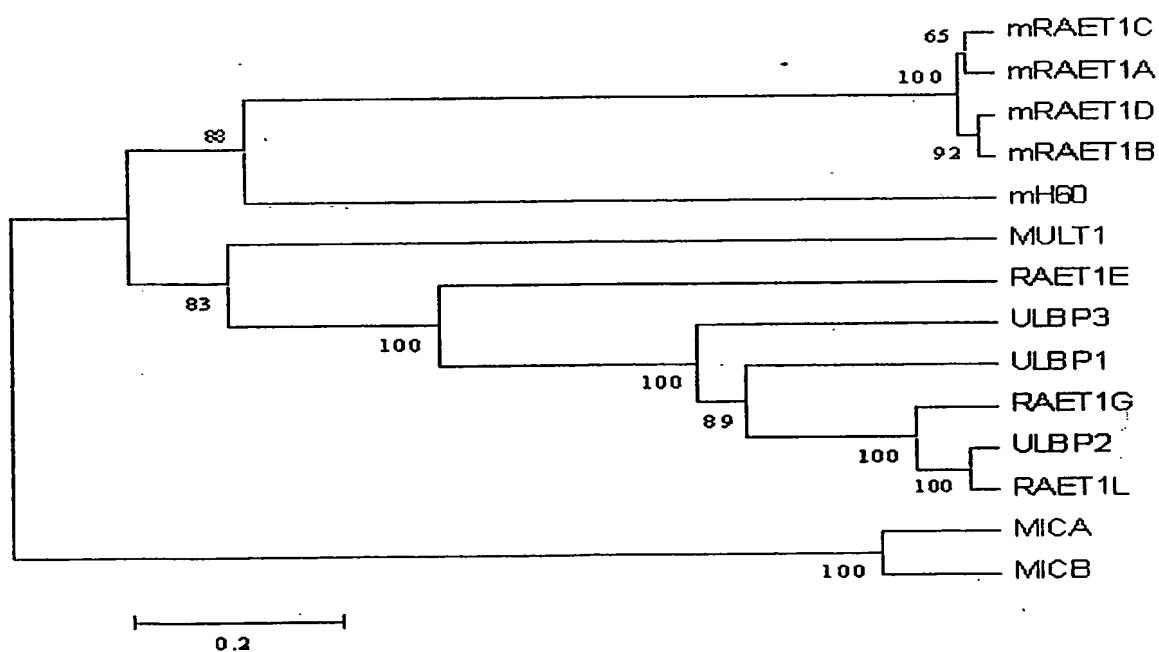


Figure 6

Raet1g
ulbp2
Raet1e
ulbp3

Raet1g
ulbp2
Raet1e
ulbp3

Raet1g
ulbp2
Raet1e
ulbp3

Raet1g
ulbp2
Raet1e
ulbp3

Raet1g

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Raet1g
ulbp2
Raet1e
ulbp3

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TELYHYDCGNKTVTPVSPLGKKLNVTAWKAQNPVLREVVVDILTEQLRDIQLENYTPKEPL 119
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TLQVEMFCQHEAERCTGASWQFTINGEKSLLFDAMNMWTVINHEASKIKETWKKDRGLE 178
TLQVRMSCECEADGYIRGSWQFSFDGRKF~~FL~~FDSSNNRKWTVHAGARRMKEKWEKDSL~~T~~ 180

[Raet1g alternative splice ending
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ulbp2
Raet1e
ulbp3

SAGGTV

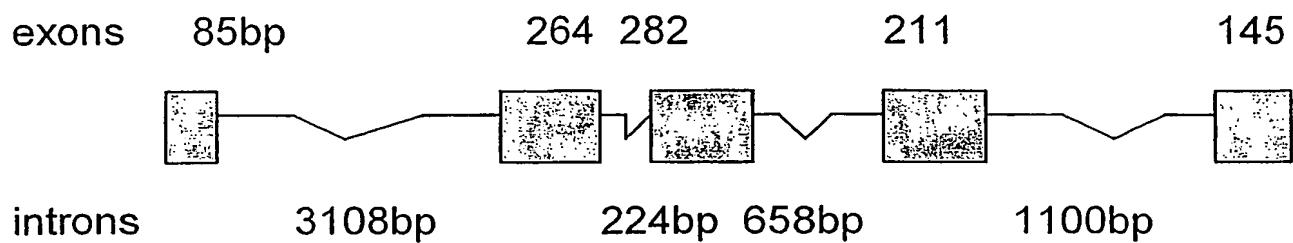
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KYFRKLSKGDCDHW~~I~~REFLGHWEAMPEPTVSPVNASDIHWS~~SS~~SLPDRW~~W~~ILGAFILLVL 238
TEFKMVS~~M~~RDC~~K~~SW~~I~~RDFLMHRKKR~~E~~PTA--PPTMAPGLAQPKAIATT~~L~~SPWSEL~~L~~IL- 237

LICSRHS~~L~~TQSHGHHPQSLQ~~OP~~PHP~~PL~~LHPTW~~L~~RLWS~~D~~SYQIAKRP~~L~~SGGHV~~T~~RV~~T~~ 297
PCFILPGI----- 246
MGIV~~L~~ICVWWONGEWQAGLW~~PL~~RTS----- 263
CFILPGI----- 244

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Figure 7

RAET1G



RAET1G2

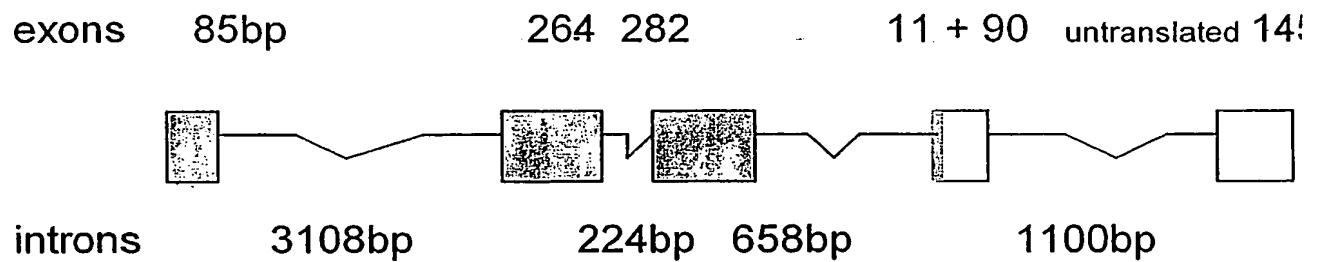


Figure 8

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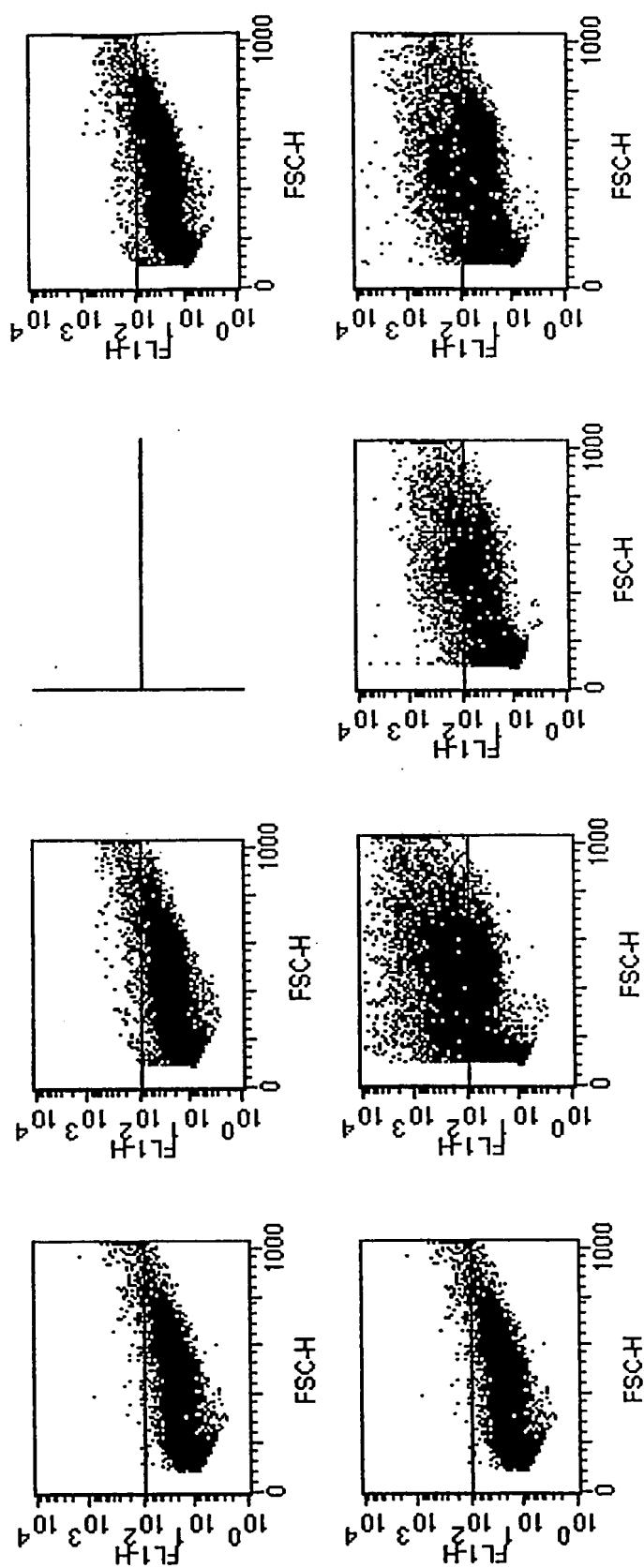


Figure 9

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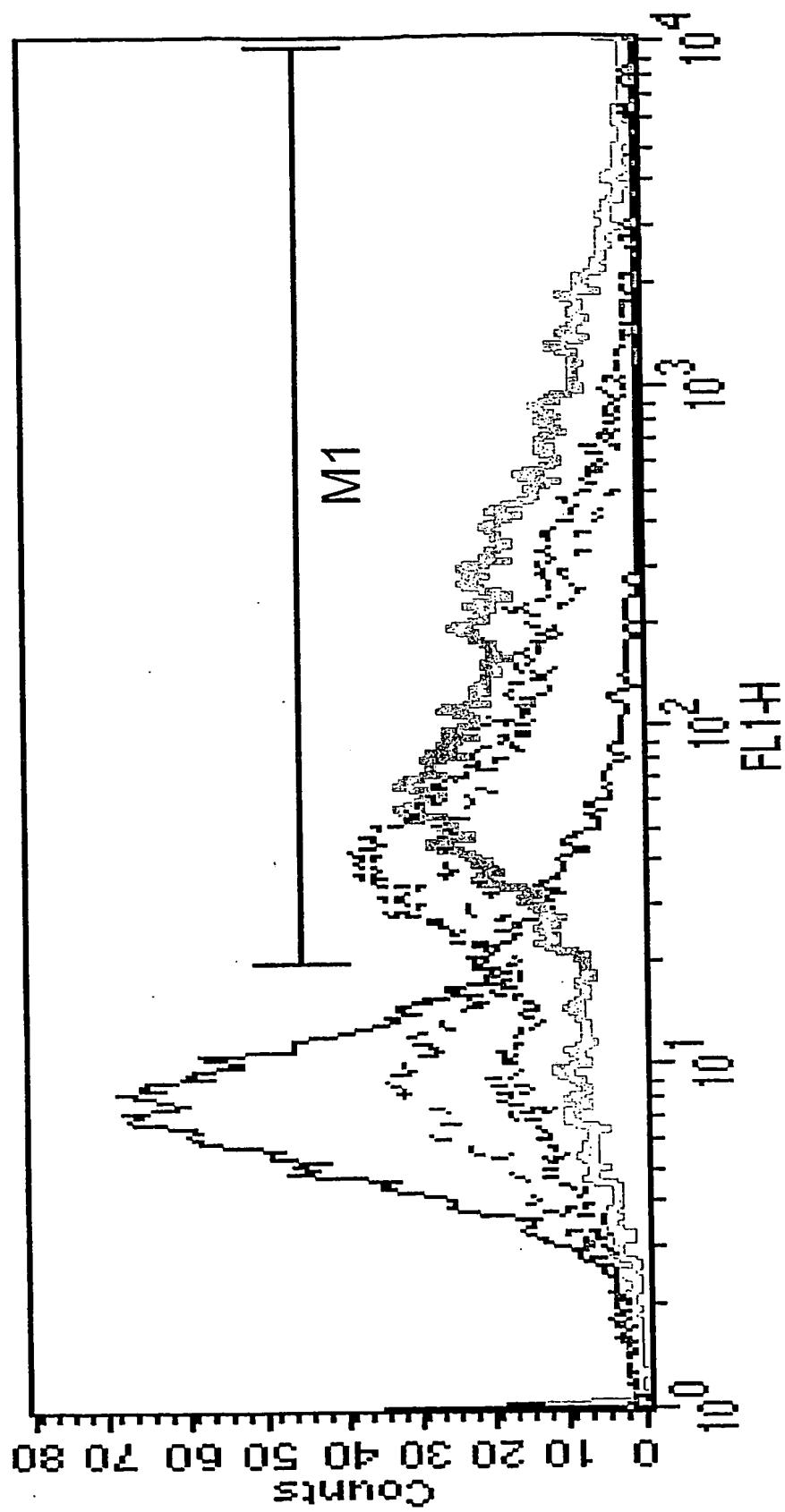


Figure 10.

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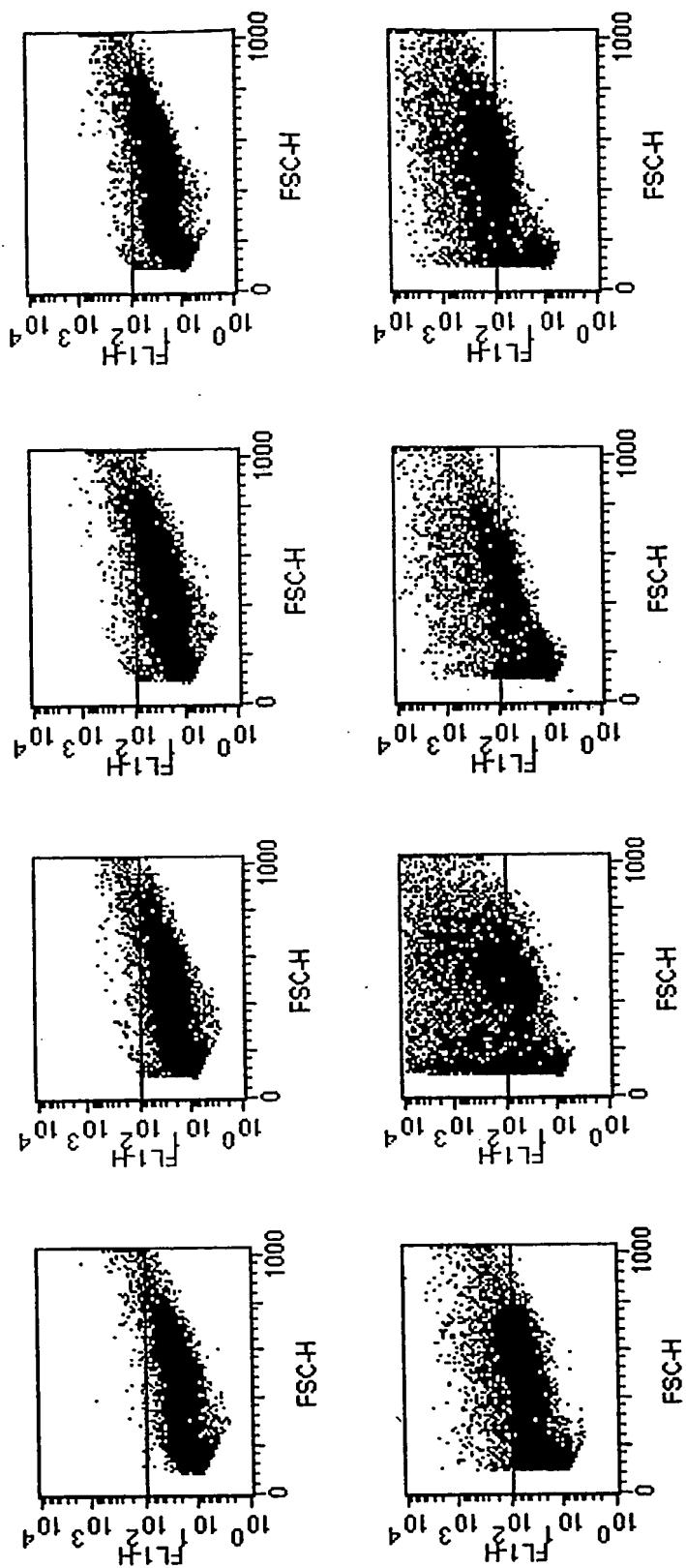


Figure 11

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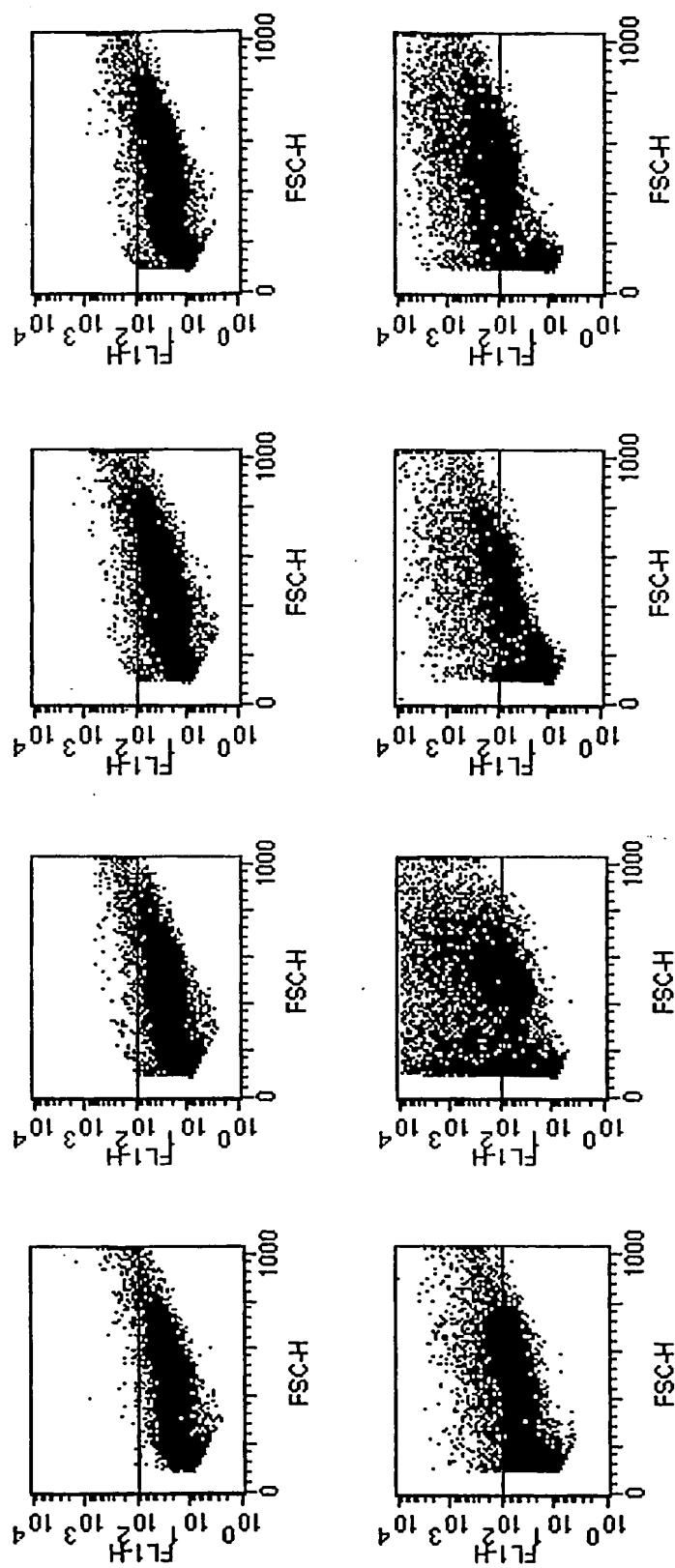


Figure 11

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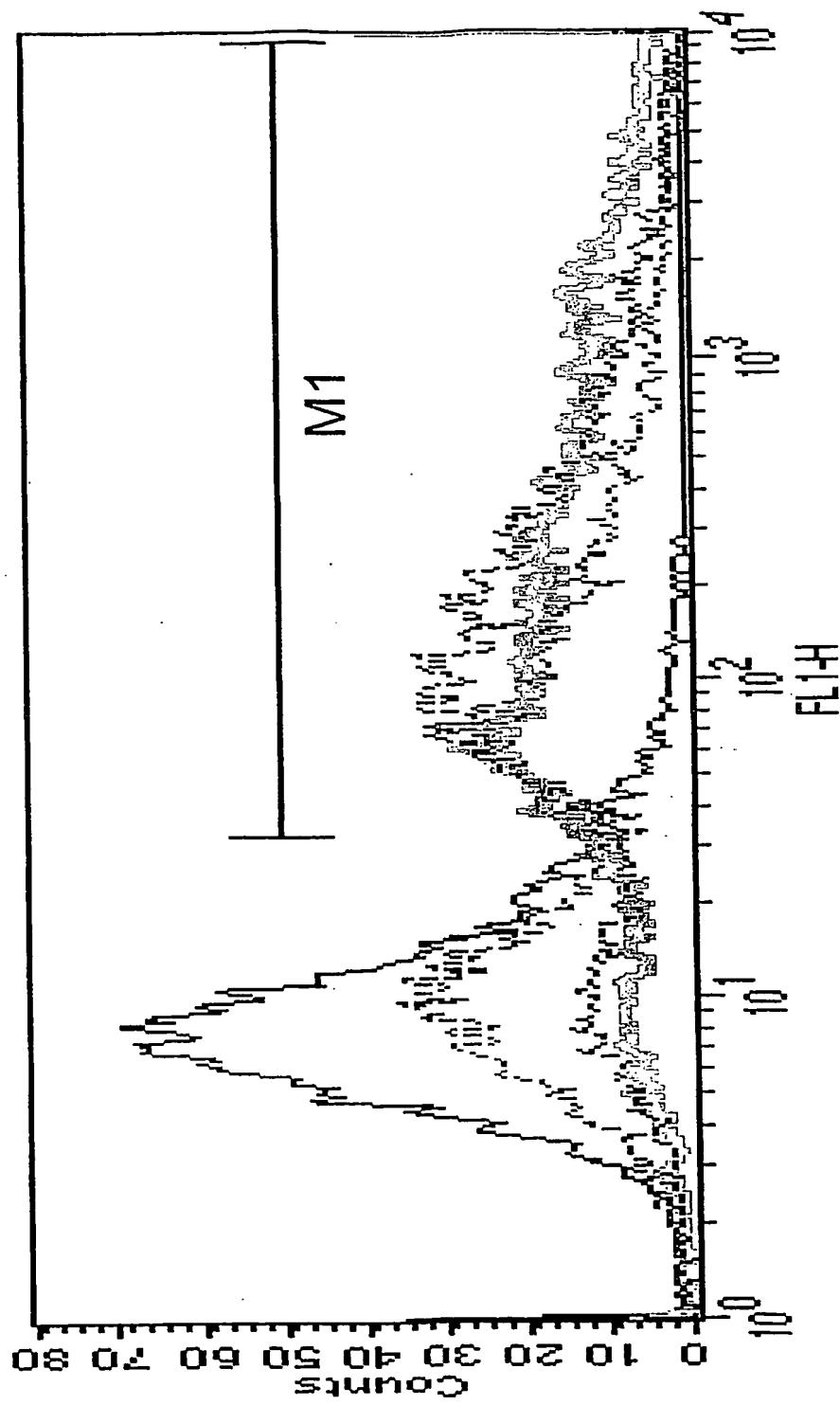


Figure 12

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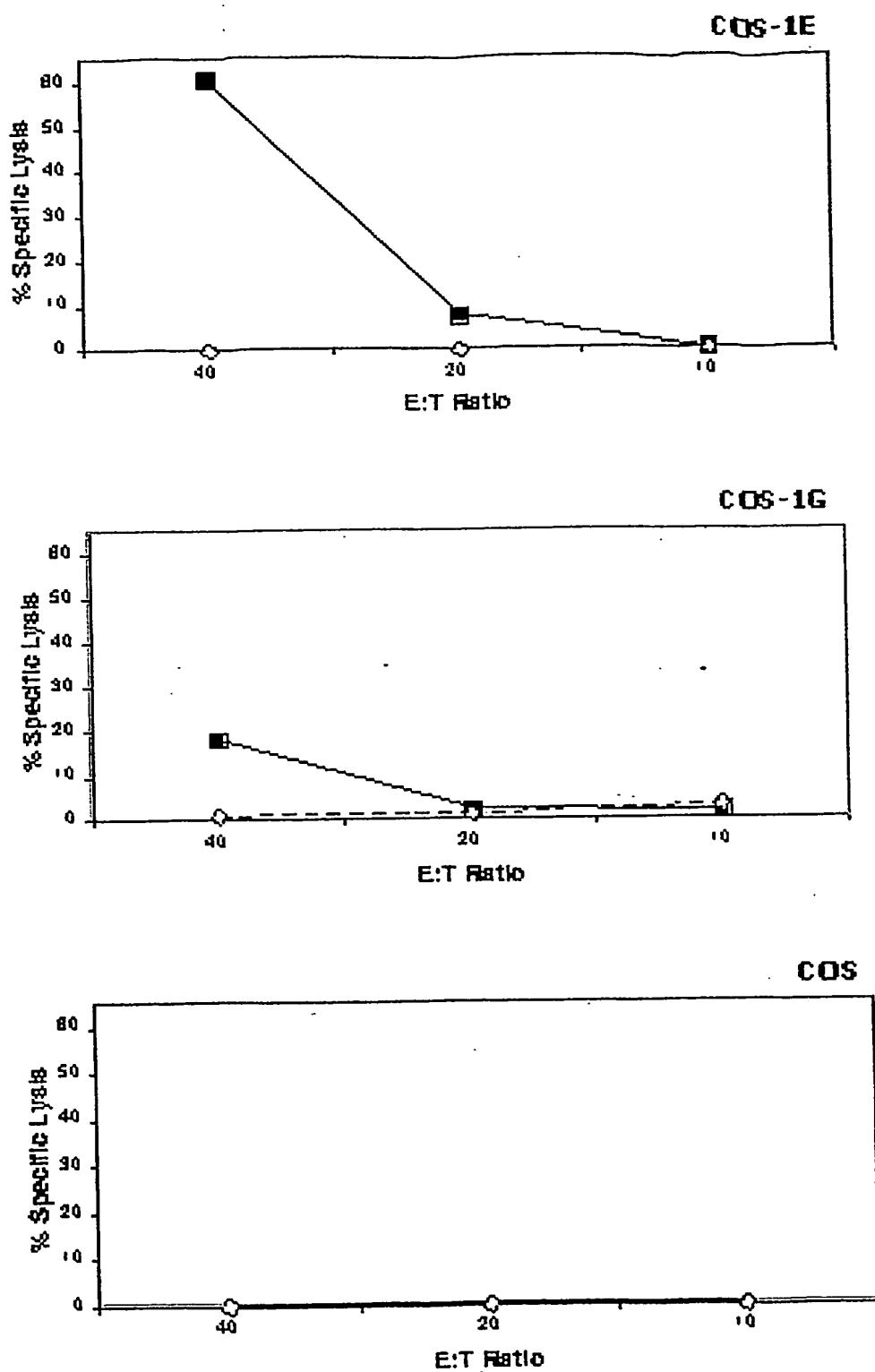


Figure 13

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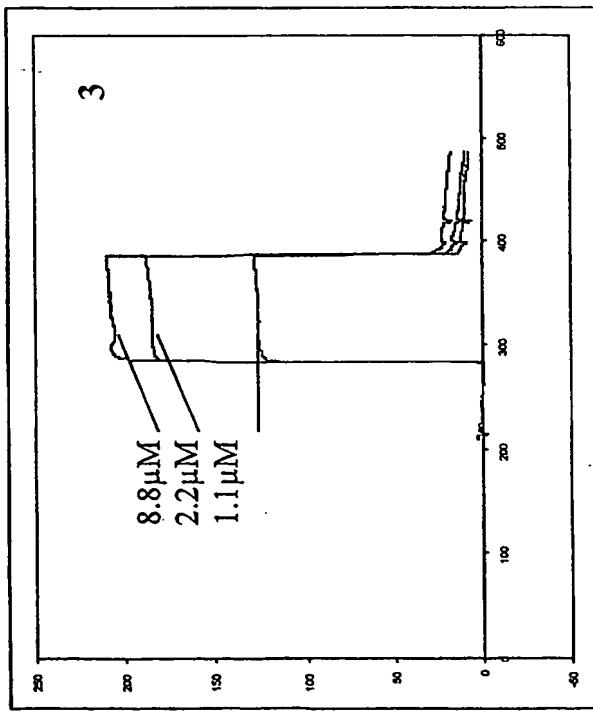
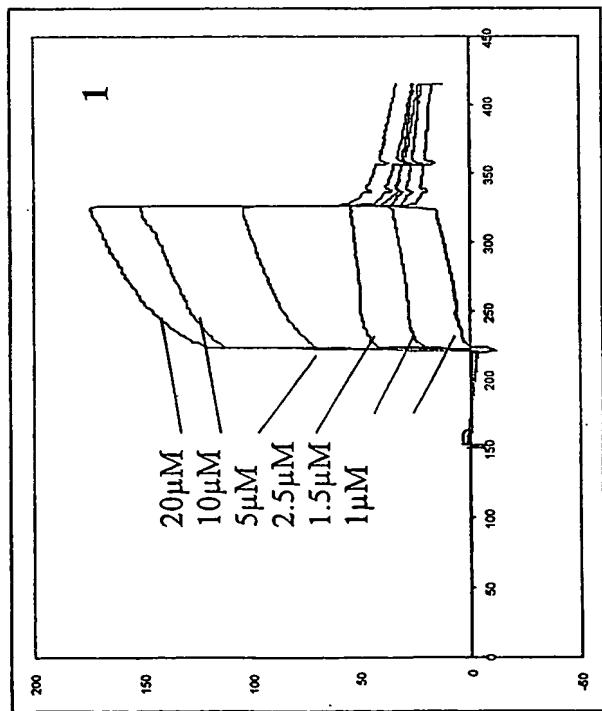
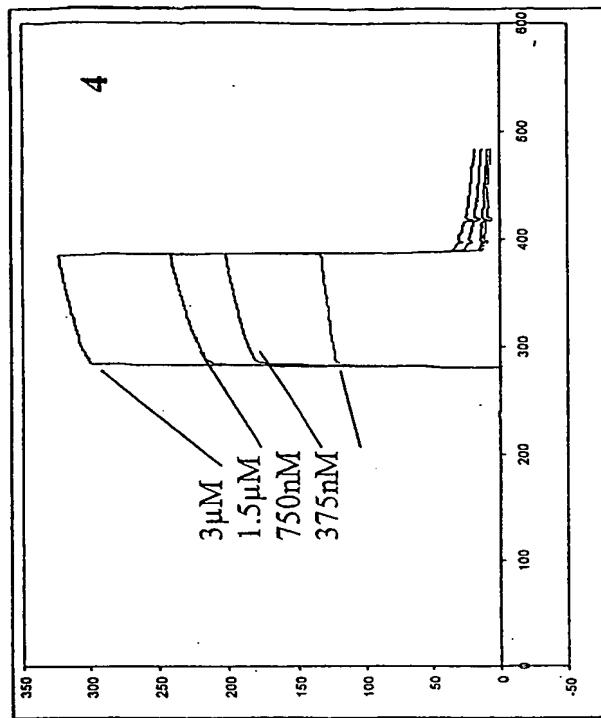
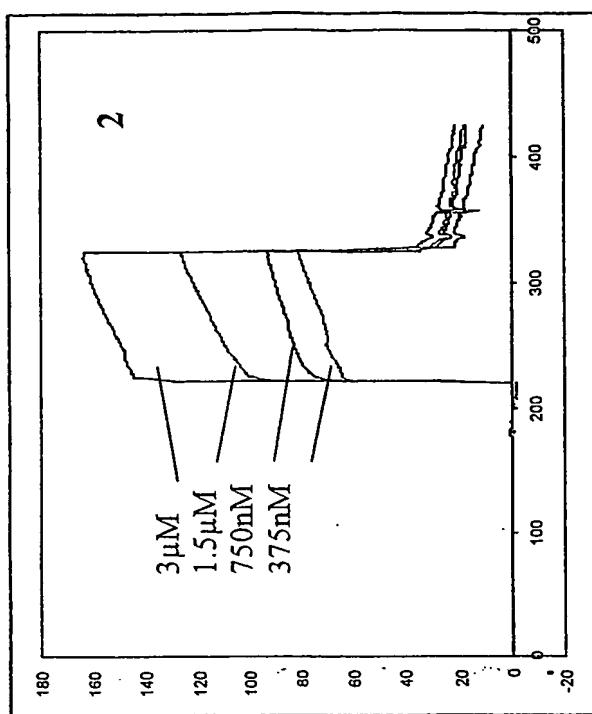


Figure 14